

■ 08-476,567



F16, 2a

60 70 TCGCCGCTCTCACTTCG <<< <<<	140 AGCACCTTCCACC	10 220 CCAGGGTGAGAAA Q G E K Q G d K	290 CCTGGACTGTGAG L D C E L X S X	60 GGGCTCCAACAGT GSNS
50 CACCAATCTCCTCGC <<< <<	90 140 120 120 140 GGCTCGTGGCCCTGTCCACCGTCCATCATCCCGCCGGCCACCGCTCAGAGCACCTTCCACC >>>> >>>>> >>>>>>>>>>>>>>>>>>>>>>>	200 STACTTGTGCCTGCC Y L C L P Y m a L P CB XI-B	240 260 270 280 290 TATATCTGGGTTGATGGTACTGGACTGCGCTGCAAAACCCGCACCCTGGACTGTGAG Y I W V D G T G E G L R C K T R T L D C E Y I W i D G T G E G L R C K T R T L X S X CB VII 30 35 40 45	350 360 ACCTTCAGTCTGAGGGC T F Q S E G
0 CTGCTTGATTCCCAC >>>>	110 CGTCCATCATCCCGC	0 190 AACATCAAGCAAATC N I K Q M 9 I K Z v	260 270 3GAGAAGGACTGCGCT G E G L R (G E G L R (340 FTTGATGGCTCTAGT FDGSS 65
10 CGAGAATGGGAGTAGAGCCGAC	100 3GCCCTGTCCACCC >>>>	170 180 CCCACTTGAACAAAA S H L N K S H L B K) 250 GGGTTGATGGTACTC W V D G T W i D G T	320 330 TACCTGAGTGGAATT L P E W N t n l z r
1 10 20 30 40 50 60 70 CCGAGCCGAGAATGGGAGTAGACTGCTTGATTCCCACCCCAATCTCCTCGCCGCTCTCACTTCG >>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>	80 CCTCGTTCTCGTGGCTCGTGGCC >>>>>>>>>	150 160 170 180 190 200 210 220 ATGGCCACCTCAGCAAGTTCCCACTTGAACAAAACATCAAGCAAATGTACTTGTGCCTGCC	230 240 GTCCAAGCCATGTATATCTG V Q A M Y I W V Q A M Y I W CB VII 25 30	300 310 320 330 340 350 360 370 CCCAAGTGTAGAAGTTACCTGAGTGGAATTTTGATGGCTCTAGTACCTTTCAGTCTGAGGGCTCCAACAGT PKCVEELPEWNFDGSSTFPQSEGSNS PKKPastnlzr 60 65 70
a: -	a: CCTCC	150 a: ATGCCCACC b: M A T c: AcA T d: CB X-D	a: GTCCAb: V Cc: V Cd: 25	300 a: CCCAA b: P K c: P K

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FIG. 2b

GACATGTATCTCAGCCCTGTTCGGGACCCCTTCCGCAGAGATCCCAACAGGTGTTCTGTGAA D M Y L S P V A M F R D P F R D P N K L V F C E M Y L V P a A M F R D P F K R D P N X L V F C E	S	GTTTTCAAGTACAACCGGAAGCCTGCAGAGCCAATTTAAGGCACTCGTGTAAACGGATAATGGACATGGTGAGC V F K Y N R R P A E T N L R H S C K R I M D M V S V F X Y N k r P A E T N L X X t C M B M V S CB XIV CB XII-C	105 110 115 120	AACCAGCACCCTGGTTTGGAATGGAAGAGTATACTCTGATGGGAACAGATGGGCACCCTTTTTGGTTGG	600 610 620 630 640 650 660 670 TCCAATGCCTATGGCCAAGGTCCGTATTACTGTGGCGCCAAAAGCCTATGGCAGGATATC S N G F P Q G P Y Y C G V G A D K A Y G R D I S N C F X G P Q a 160 165 170
380 a: GACATGTATCTC b: D M Y L c: M Y L		a: GTTTTCAAGTAC b: V F K Y c: V F X Y d:	100	a: AACCAGCACCCC b: N Q H P c: N Q X P d:	600 a: TCCAATGGCTTT b: S N G F c: S N C F

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FIG. 2c

680 690 710 720 730 740 GTGGAGGCTCACTACCGCGCCTGCTTGTATGCTGGGGTCAAGATTACAGGAACAAATGCTGAGGTCATGCCTGCC	750 760 770 780 790 800 810 820 CAGTGGGAATCCTGTGGAATCCGCATGGGAGATCATCTCTGGGTGGCCCGTTTCATCTTG WE F Q I G P C E G I R M G D H L W V A R F I L Q W E F Q I G P C E G I d M 215 220	830 840 860 870 880 890 CATCGAGTATGTGAACTTTGGGGTAATGGTGCAACCTTTGACCCCAAGCCCATTCCTGGGAACTGGAATGGTGCA H R V C E D F G V I A T F D P K P I P G N W N G A 225 240 245	900 910 920 930 940 950 960 970 GGCTGCCATACCAACTTTAGCACCATGCGGGAGGAGGAGGAGTCTGAAGCACATCGAGGAGGCCATCGAG G C H T N F S T K A M R E E N G L K H I E E A I E M X E E N G L K Y I E E A I E CB III-C 250 265 270
730 AATGCTGAG N A E N X X :-D 195	B1. TGGGTGGCC W V A 220	880 CCTGGGAAC P G N 245	96 CACATCGAG H I E Y I E 270
720 TACAGGAACA T G T 9 G T - CB VI 190	800 AGATCATCTC D H L 215	870 CAAGCCCATT K P I 240	950 G L K G L K G L K
710 3GGTCAAGAT' 3 V K I 3 i K	790 rccgcatgggi I R M G I d M	860 CCTTTGACCC	940 GGAGGAGAA7 REEN KEEN
700 TGTATGCTG(L Y A (L Y A (780 GTGAAGGAA C E G 210	850 TAATAGCAA(V I A 7	930 AGGCCATGCC K A M I M 2 CI
690 CGCGCCTGCT R A C A C T VII-G	770 ATAGGACCCT I G P I G P 1 G P	840 SACTTTGGGG D F G	920 TTTAGCACCA F S T 255
680 SCTCACTACO A H Y	760 GAATTCCAAI E F Q E F Q	830 GTATGTGAAC V C E	910 CATACCAACT H T N
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FIG 2c

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990 1000 1010 1020 1030 1040 GCGGCACCGGTACCATTCGAGCCTACGATCCCAAGGGGGCCTGGACAATGCCCGTGGTCTG RHRYHIRAYDPKGGLDNARGLTCTG Sninyq AYBPK TIX-B-1 280 285	1050 1060 1070 1080 1090 1100 1110 1120 ACTGGGTTCCACGAAACGTCCAACTCAACGACTTTTCTGCTGGTGTCGCCAATCGCAGCATCCGCATT T G F H E T S N I N D F S A G V A N R S A S I R I T S N I N Y Q G A S I R I R I C-CB III-C) 300 315 310	1130 1140 1150 1160 1170 1180 1190 CCCCGGACTGCCGCCAGGAAGAAAGGTTACTTTGAAGACCGCCCCCTCTGCCAATTGTGACCCCTTTGCA PRTVGQEKKGYFEDRRFFA	1200
1040 CCCGTG	ATCC I I -1	1190 CCCCT	AACT N *
10 BA	1110 CGCAGTGCCAGCAT R S A S I 9 A S I T IX-C-1	1) GACC	0 AAAA K K
CAAT N 295	1111 FGCCC A A T I	rrgT C 345	1260 ATACAA Y K Y K 370
1030 TGGA(L D	CAG' S 9	1180 CCAA: A N	CCA!
i i i I	OO ATCG	1 TGC	CTT
9999	110 CCAA	CCTC	1250 AGCCC' 3 P
1020 AAGG(K (K	50 1070 1080 1090 1100 SAAACGTCCAACATCAACGACTTTTCTGCTGGTGTCGCCAATC E T S N I N D F S A G V A N T S N I N Y Q (-CB III-C) 305 315	1170 CGCCC R 1	GACG1 D 1 B 2 365
1 3 3 3 4 1 2	oo GG G	1 SGCC R 3	3 3 3
SATC D B K-B-	109 3CTG A	засс D	1240 ACTGG(T G T G
1010 CCTACGATCC A Y D P A Y B P T IX-B-1	TCT(1160 TGAA(GAGI E Z
1 A A A	1080 GACTTT D F Y q 310	TIL	AAT N N
rcga R 285	108 CGAC D D Y)	гтас [,] Ү 335	1230 FCTCAJ L 1 L 1
1000 ACAT H I G	CAA(N N-C	1150 AAGG' K G	TGCCTT C L C L T VI-K
CCCA H H	0 CAT I I	I GAA. K	O ATG
SGTP X Y 1 Y	1070 CCAACATCAAC S N I N S N I N (-CB III-C)	AGAA E K	1220 3CACA'
990 CACC H 1 i 1	ACGTOTO TO STATE S	1140 CAGG2 Q 1	GTCCC V J
)GGC R n 2	50 3AAA E 3	1 3 3	I I 3
AGCAAGCGG S K R St Ks n <-B-2	106 CACC H	STCG V	1210 3CCAT A I
980 AAACTAAGCAA(TH F	1130 3GACT(1 T	GAA(E
CTA L L T I	SO FIGGGTTC G F	1) CCCCGG/ P R P R T IX-E 325	3ACA T
AAAO K X Z	105 ACT T T 300	CCC P P P P P 325	1200 GTGA(V '
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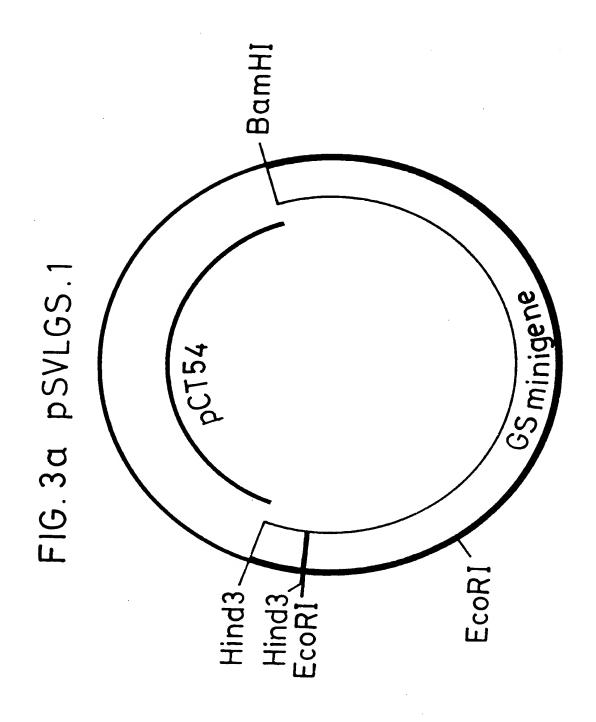
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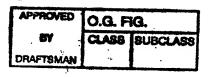
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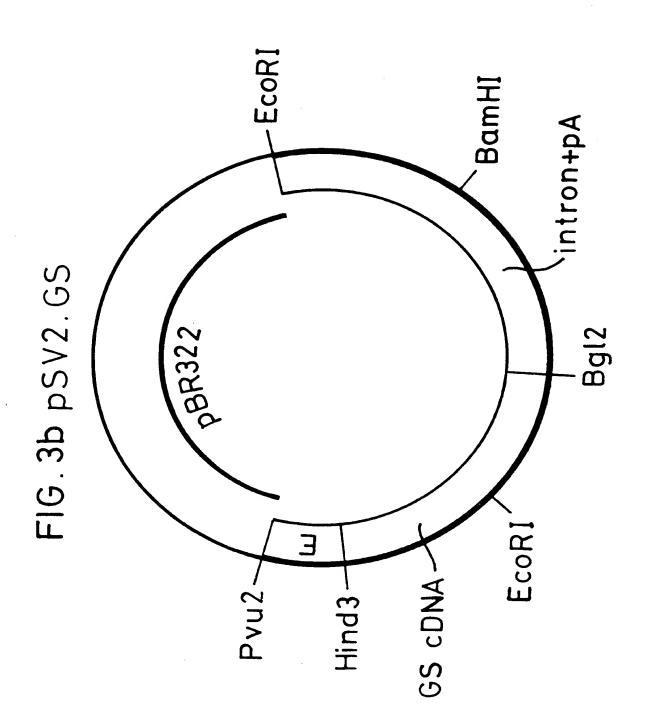
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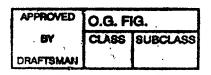
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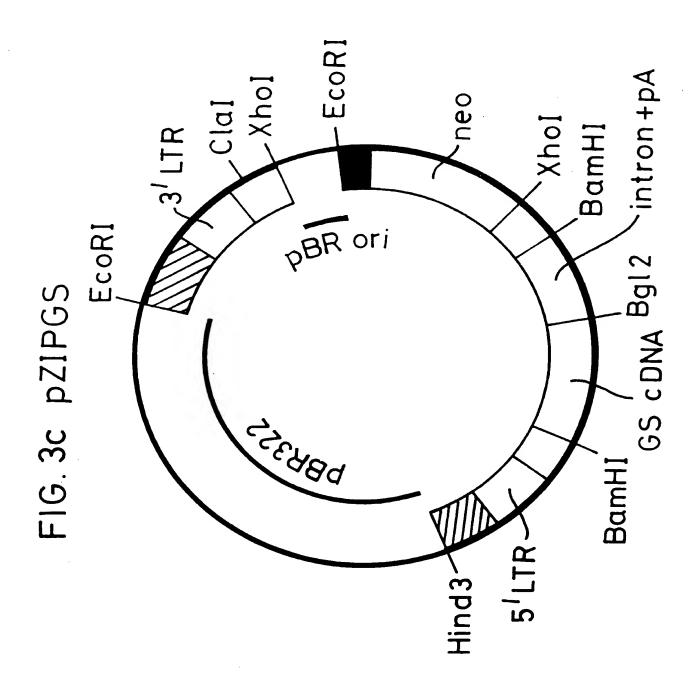




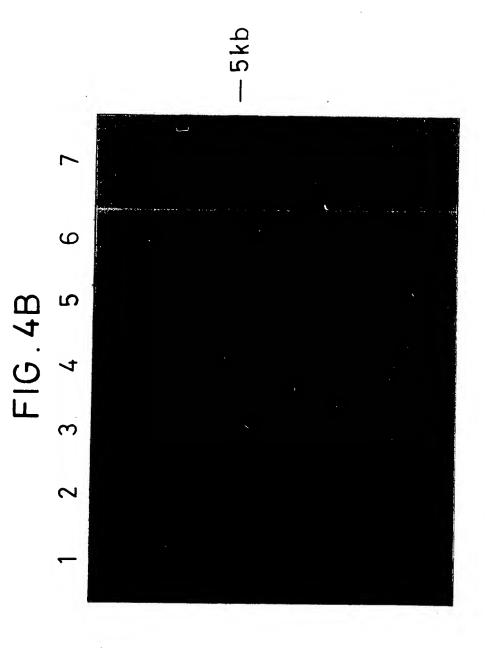


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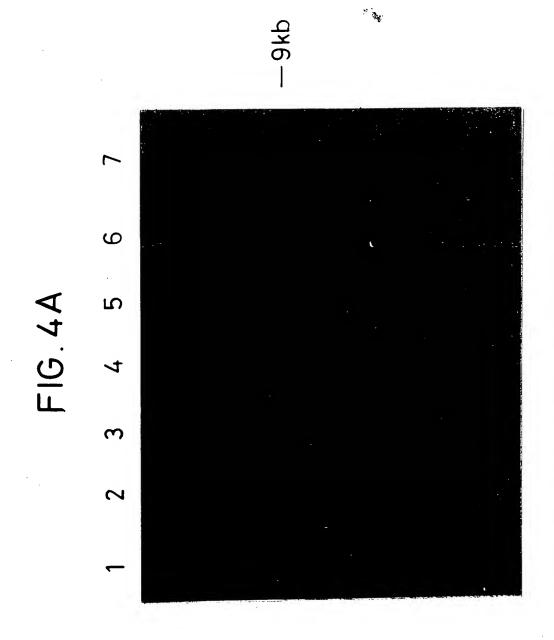


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APPROVED O.G. FIG.
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FIG.5

